Amendment to the Claims:

Claims 1-37 (Canceled)

- 38. (Currently amended) A transgenic mouse whose genome comprises a <u>null homozygous</u> disruption in an MC3-R gene-allele; said allele comprising the sequence of set forth in SEQ ID NO:1; said null allele comprising exogenous DNA, said exogenous DNA comprising a gene encoding a visible marker, wherein in a male transgenic mouse said gene is capable of expression in the testis, wherein as a result of the disruption, the transgenic mouse exhibits, relative to a wild-type mouse, passive behavior or a decrease in attempts to escape.
- 39. (Currently amended) The transgenic mouse of claim 38, wherein the disruption was produced using a targeting construct comprising the sequences set forth in SEQ ID NO:3 and SEQ ID NO:4).

Claim 40 (canceled)

- 41. (New) The transgenic mouse of claim 38 wherein said mouse is heterozygous for said disruption.
- 42. (New) The transgenic mouse of claim 38 wherein said mouse is homozygous for said disruption.
- 43. (New) The transgenic mouse of claim 42 wherein said mouse exhibits, relative to a wild-type control mouse, passive behavior.
- 44. (New) The transgenic mouse of claim 38 wherein said exogenous DNA further comprises a gene encoding a selection marker.
- 45. (New) The transgenic mouse of claim 44 wherein said gene is a neomycin resistant gene.
- 46. (New) The transgenic mouse of claim 38 wherein said visible marker is lacZ.
- 47. (New) A method of identifying an agent capable of modulating activity of a MC3-R gene or MC3-R gene expression product, the method comprising:
 - a. administering a putative agent to the transgenic mouse of claim 38;
 - b. administering the agent to a wild-type control mouse; and
 - c. comparing a physiological response of the transgenic mouse with that of the control mouse, wherein said physiological response is a change in passive behavior;

d. wherein a difference in the physiological response between the transgenic mouse and the control mouse is an indication that the agent is capable of modulating activity of the MC3-R gene or MC3-R gene expression product.